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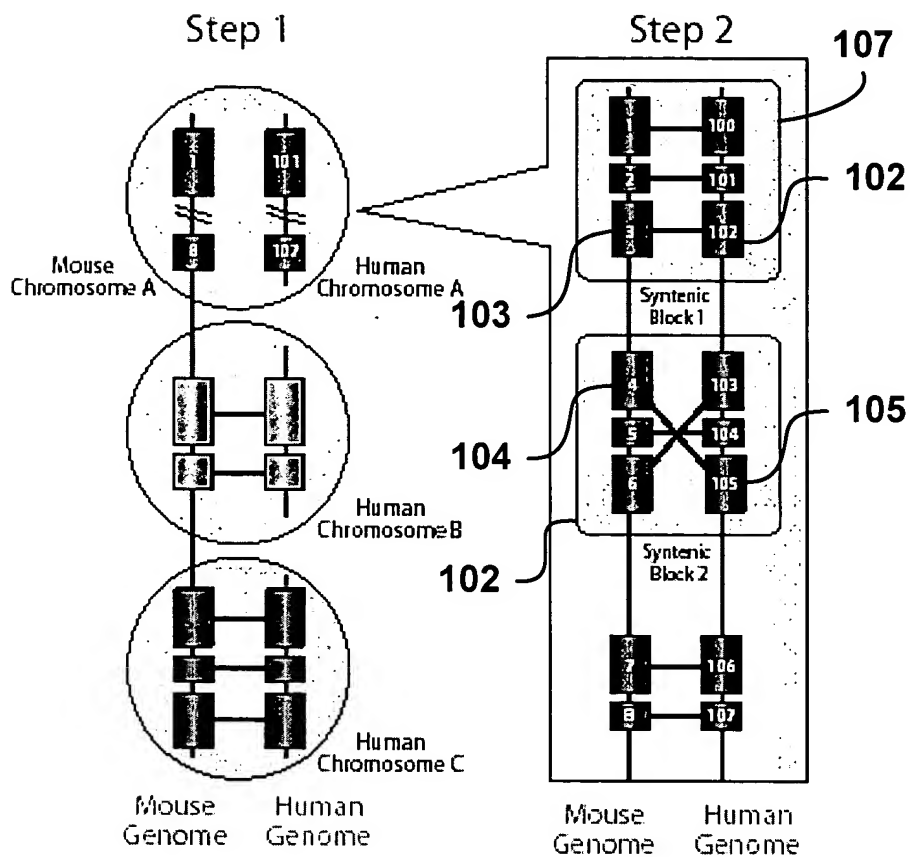


Figure - 1

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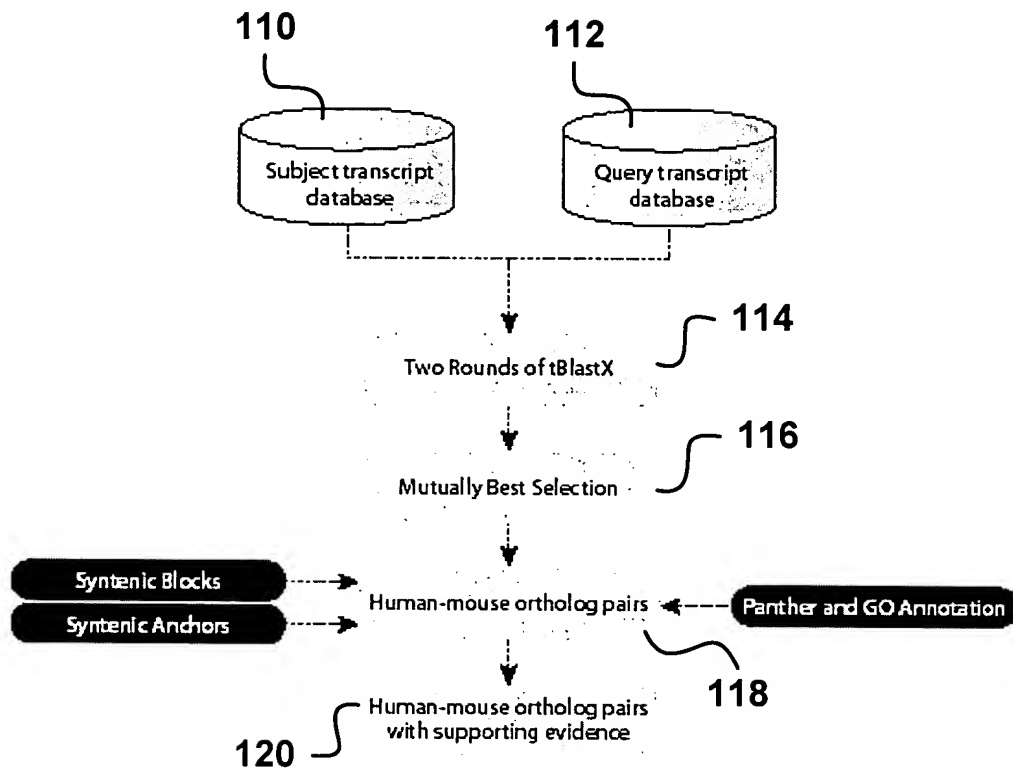


Figure - 2

Title: METHOD FOR IDENTIFYING, VIEWING, AND ANALYZING SYNTENIC AND
ORTHOLOGOUS GENOMIC REGIONS BETWEEN TWO OR MORE SPECIES

Inventors: Moore, et al

Atty. Ref.: 9692-000030

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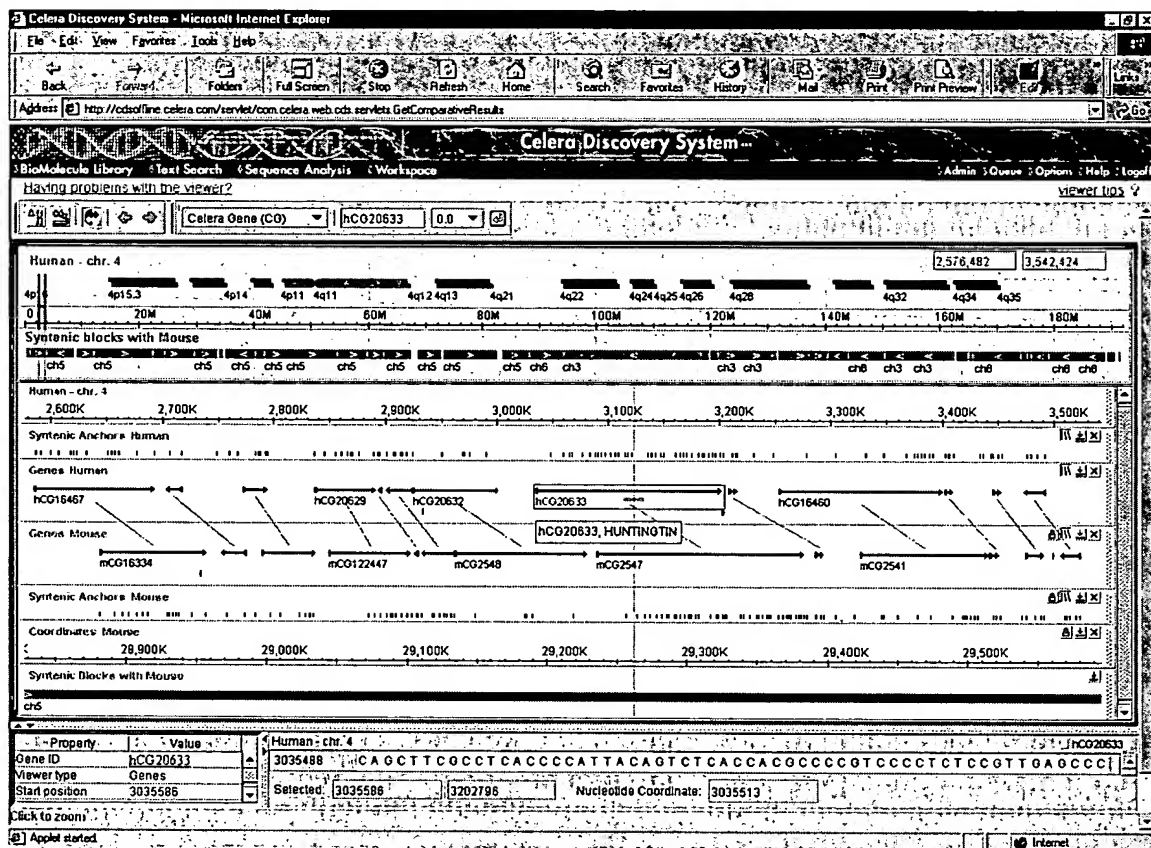


Figure - 3A

Inventors: Moore, et al
Atty. Ref.: 9692-000030

[illegible]

Figure - 3B

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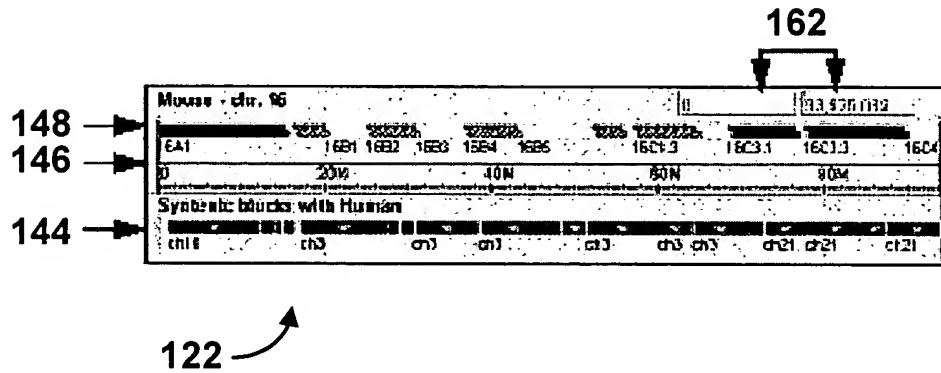


Figure - 5

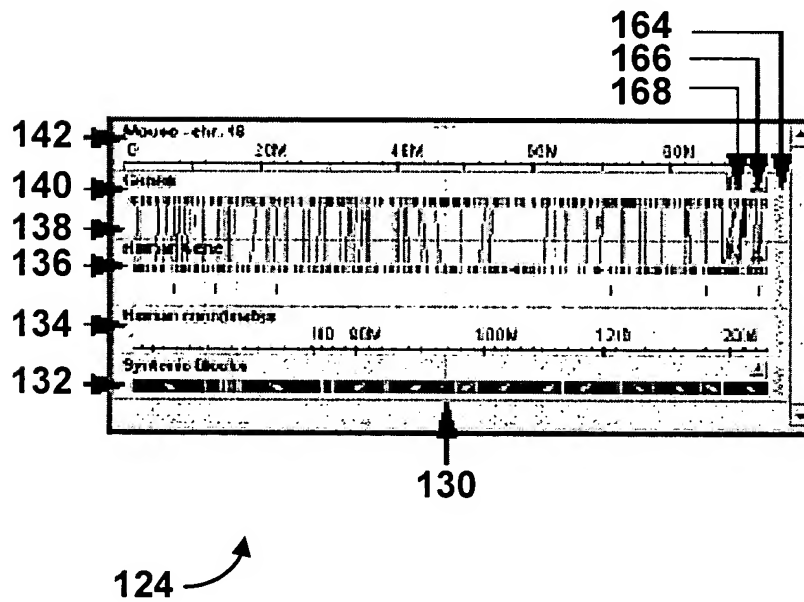


Figure - 6

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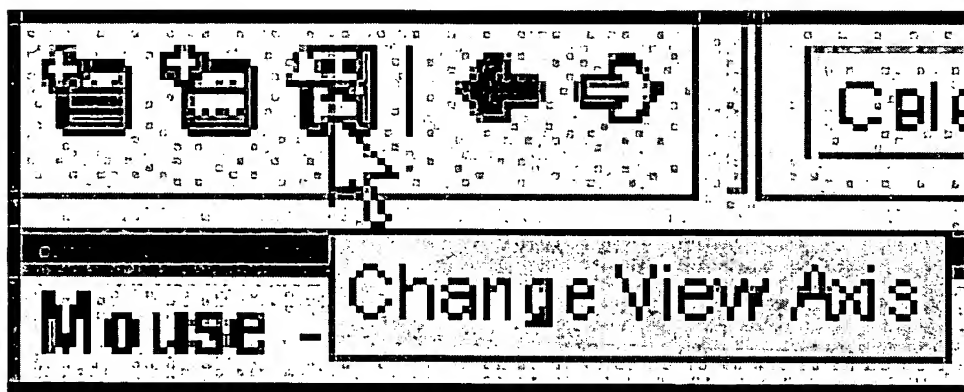


Figure - 7

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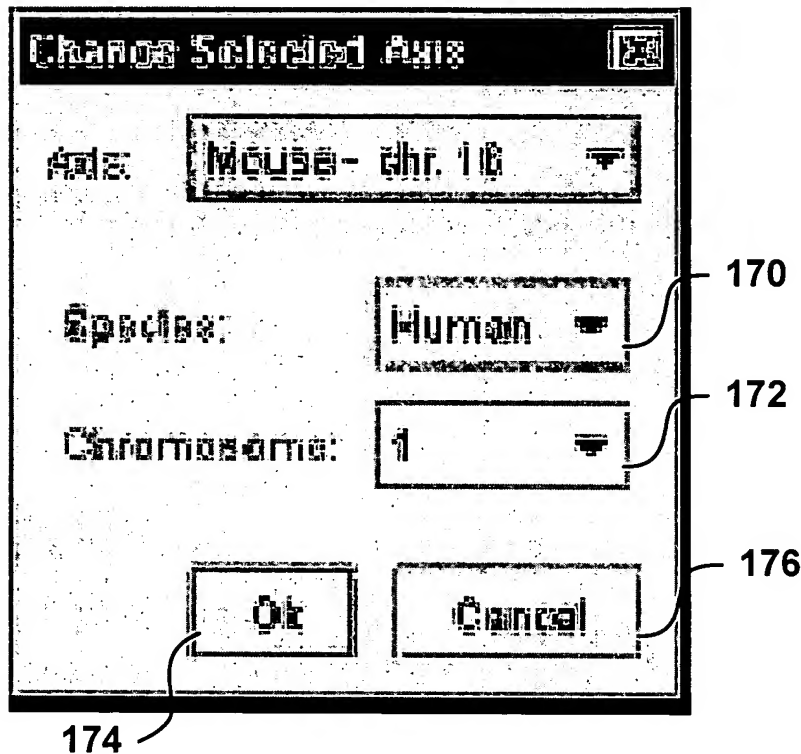


Figure - 8

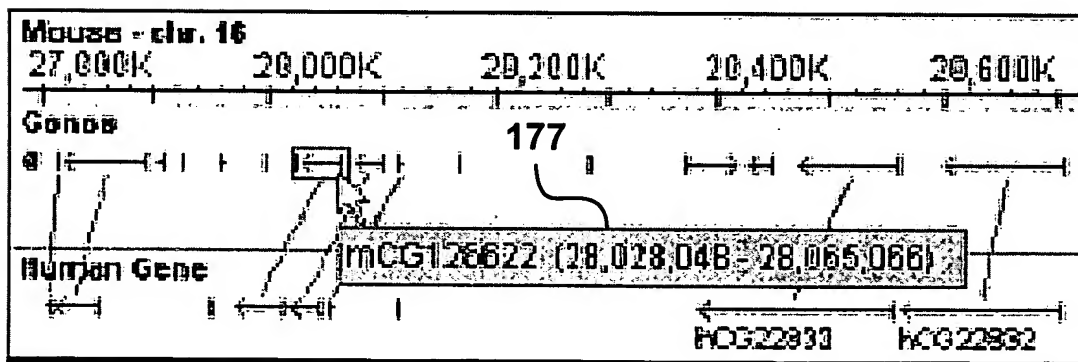


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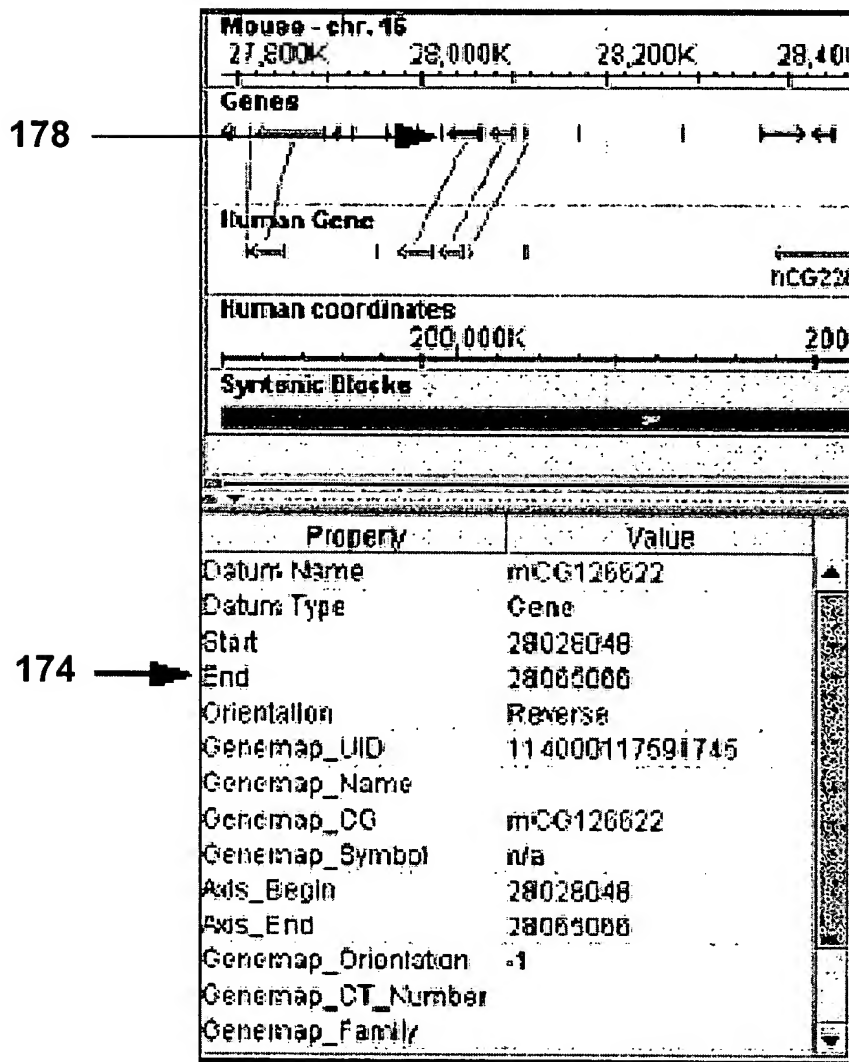


Figure - 10

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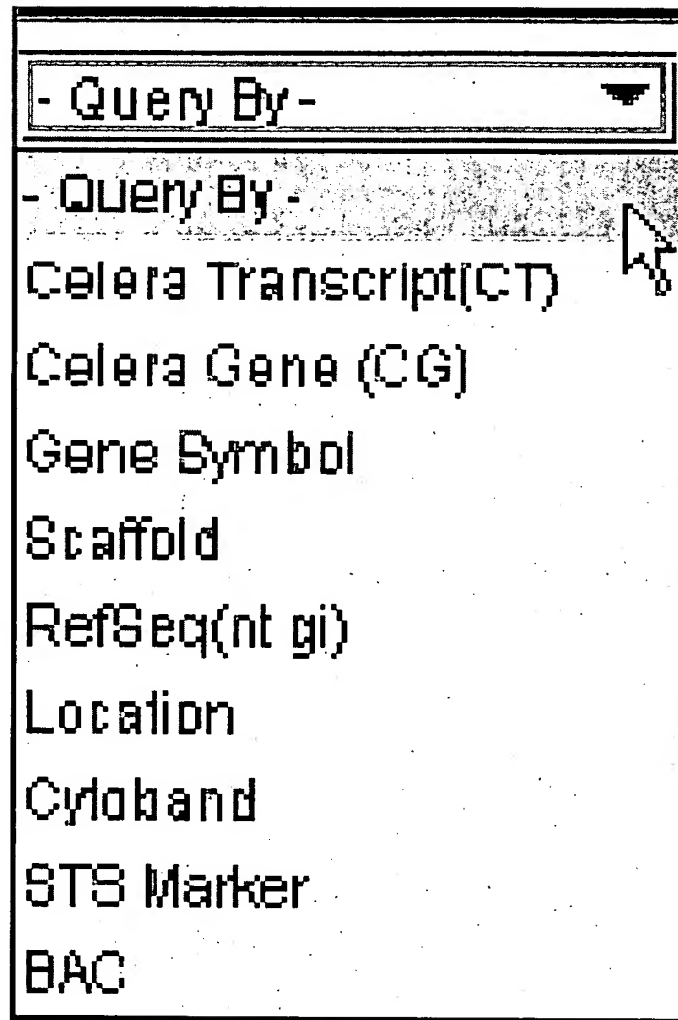


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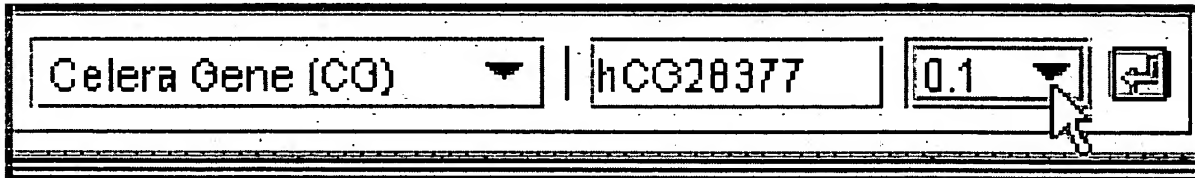


Figure - 12

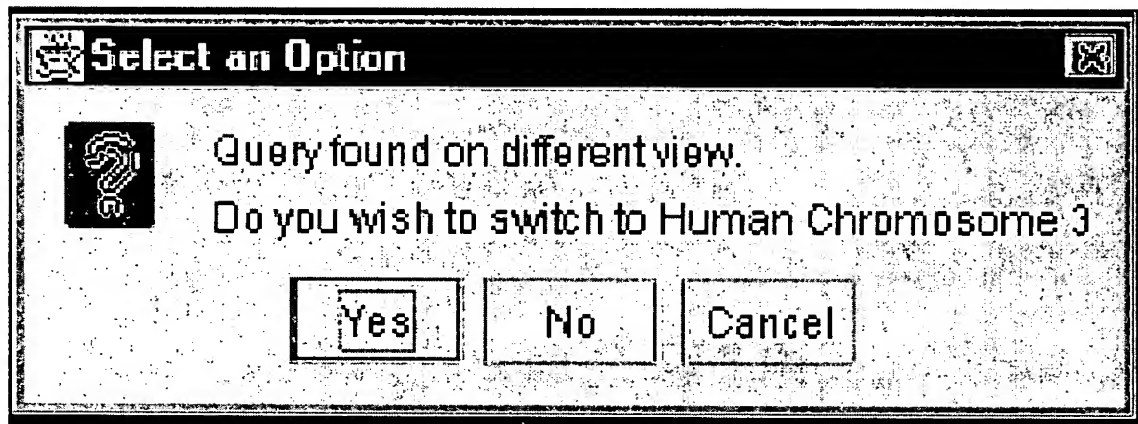


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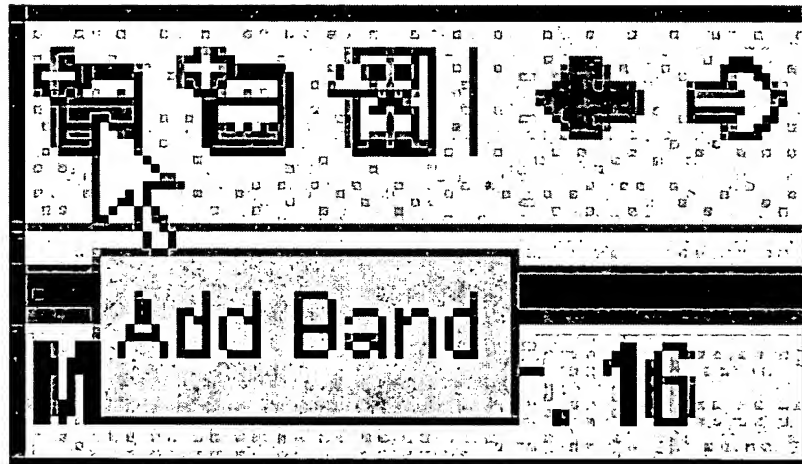


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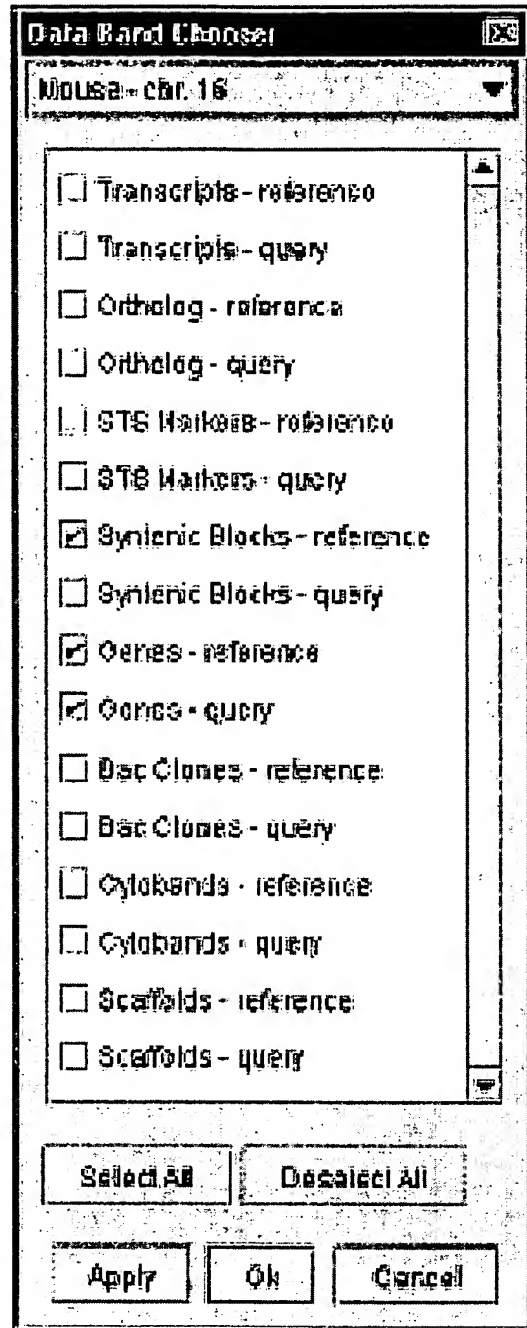


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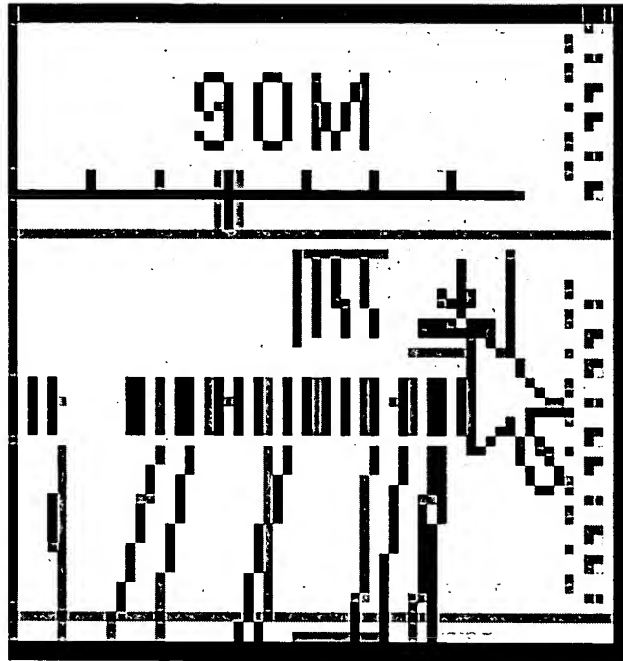


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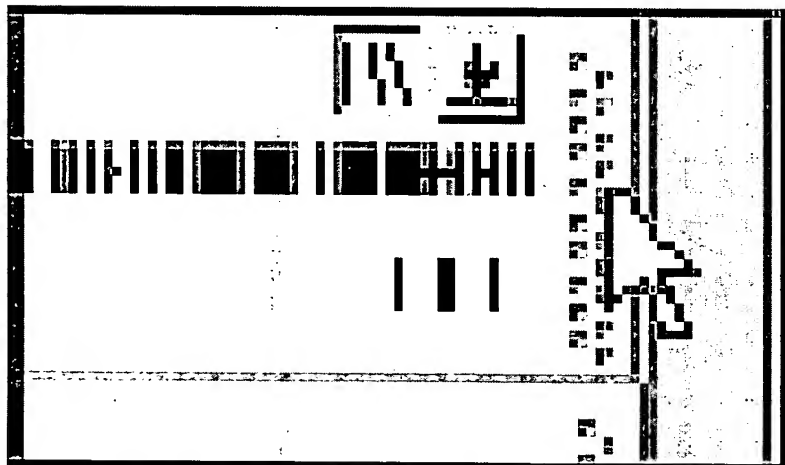


Figure - 17

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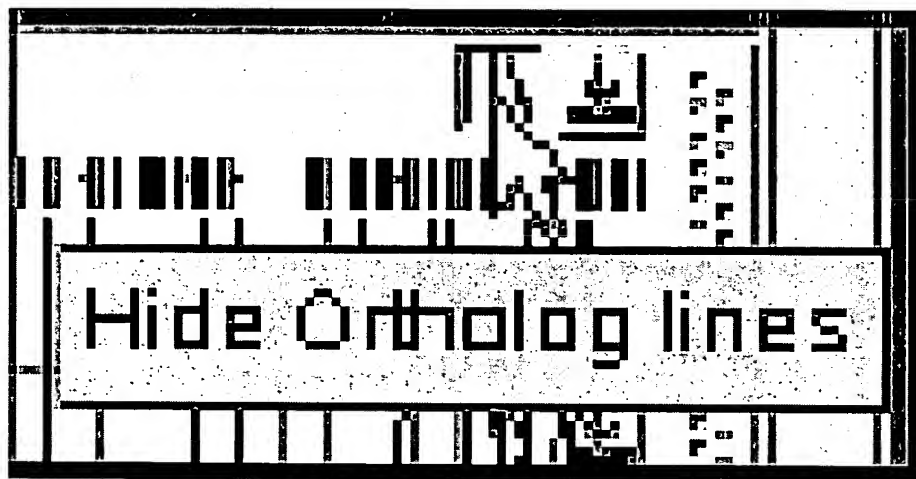


Figure - 18

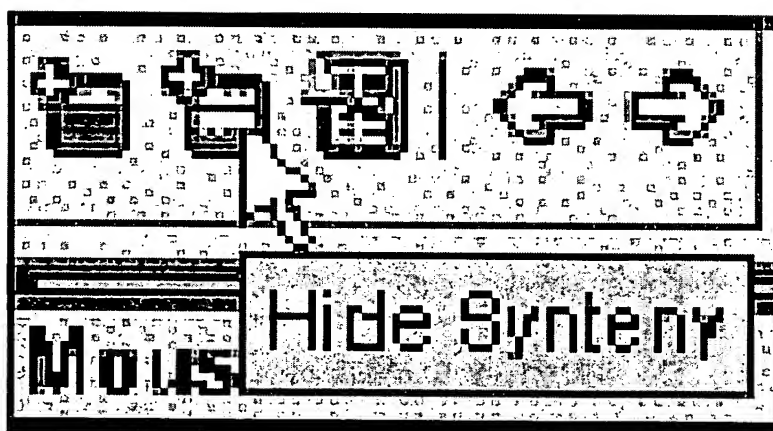


Figure - 19

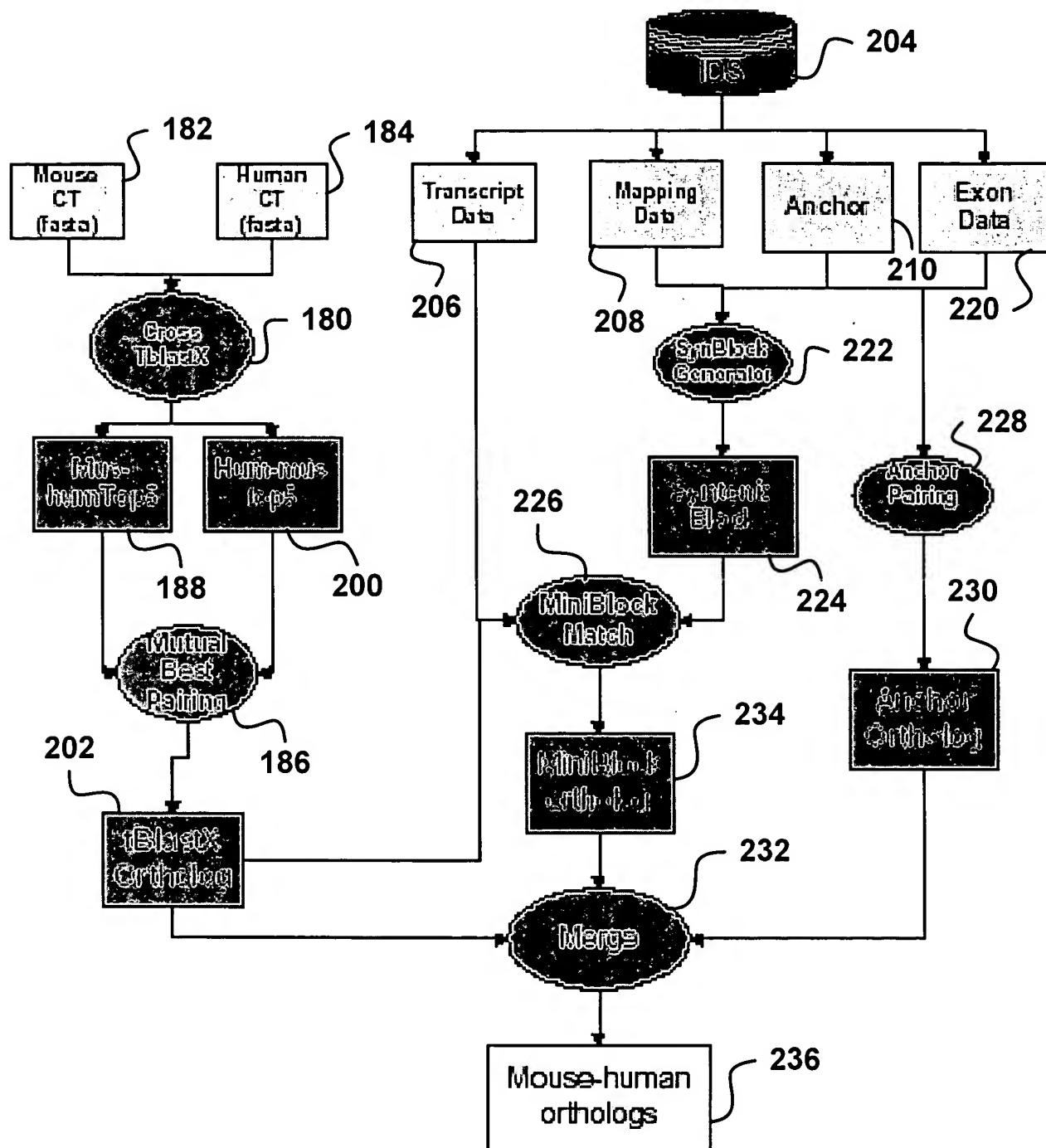


Figure - 20

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Inventors: Moore, et al

Atty. Ref.: 9692-000030

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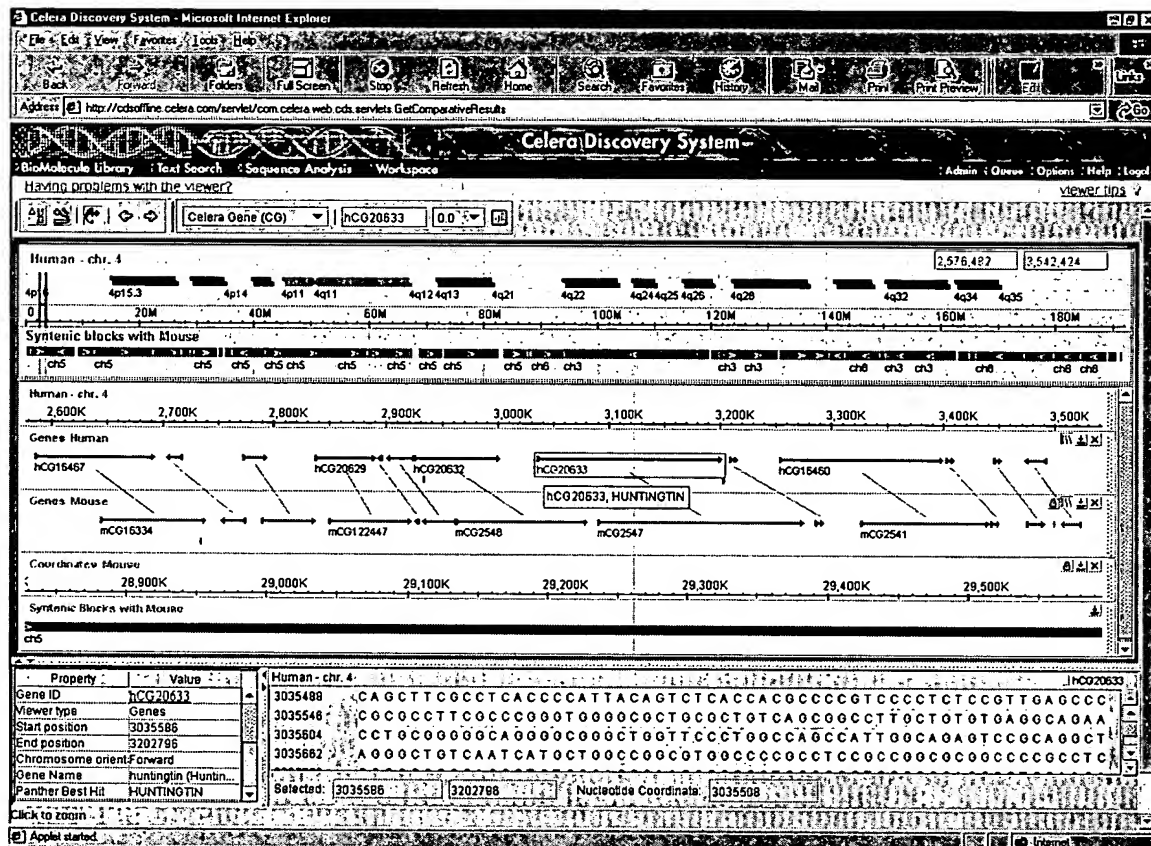


Figure - 21

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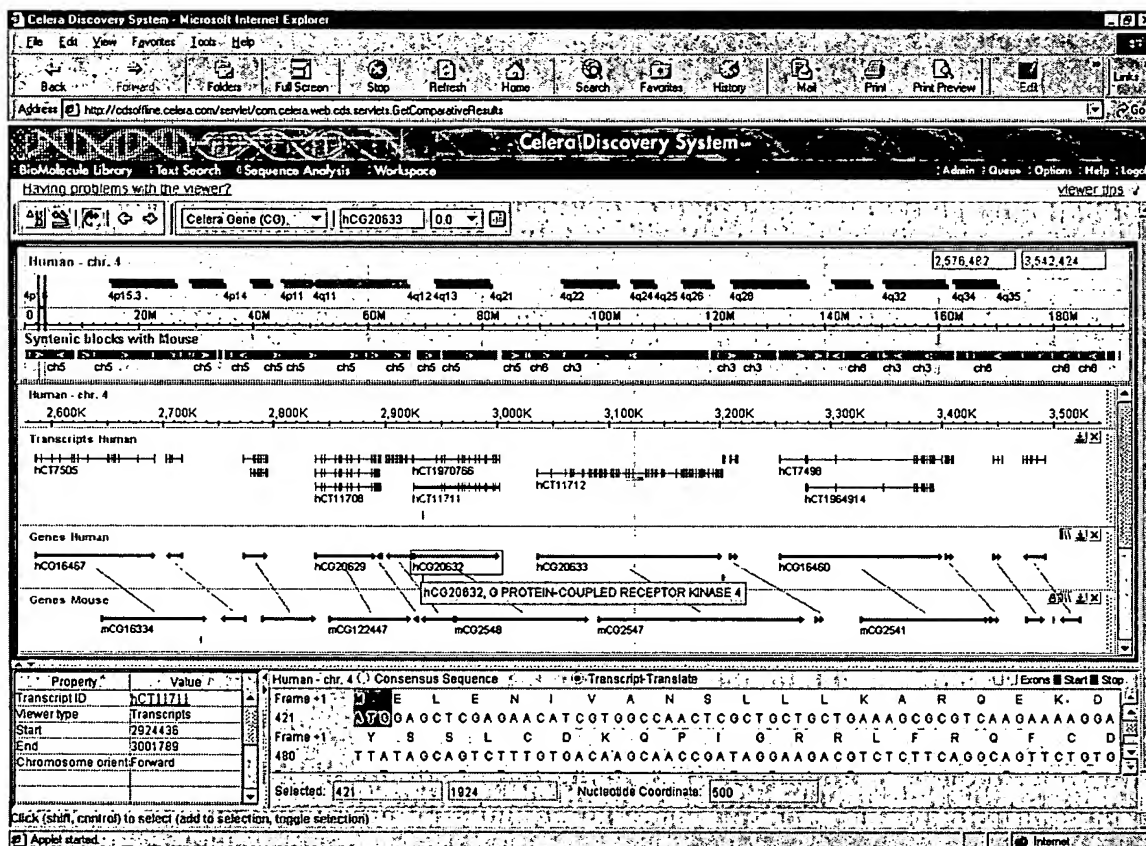


Figure - 22

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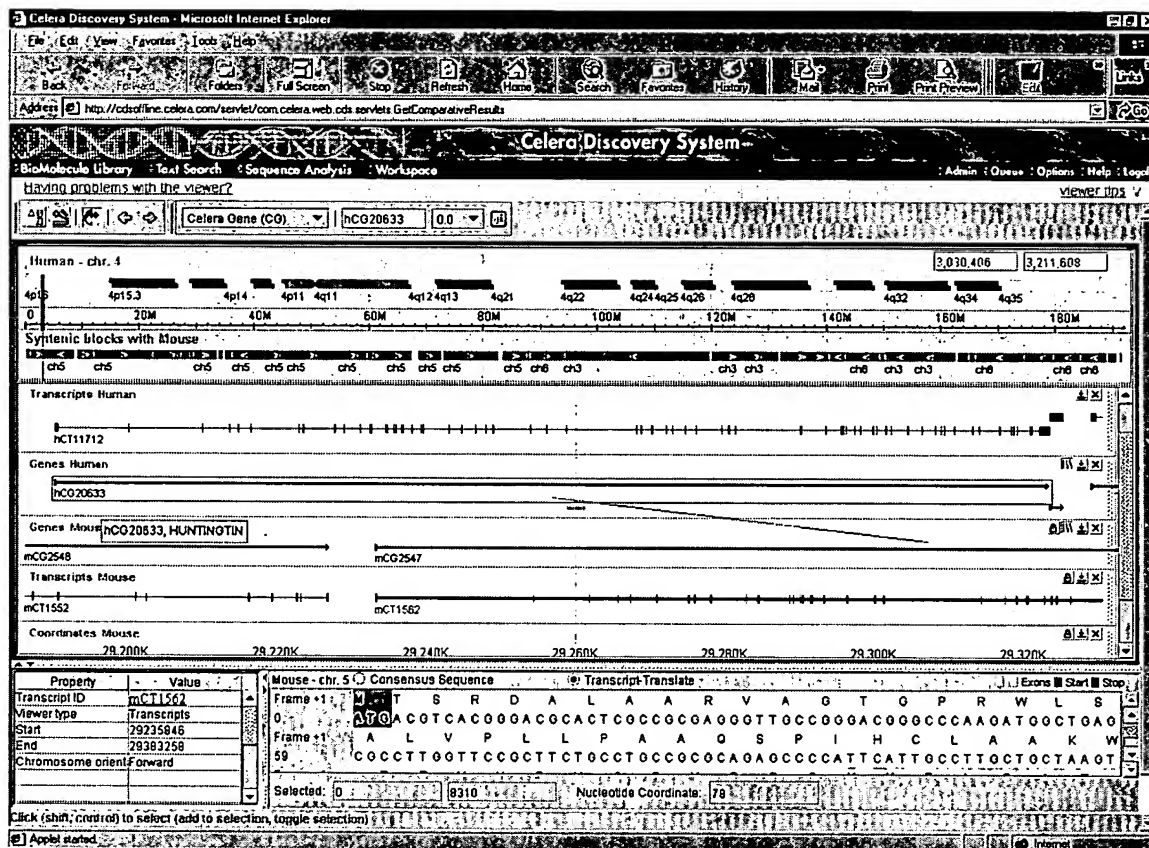


Figure - 23

Inventors: Moore, et al
Atty. Ref.: 9692-000030

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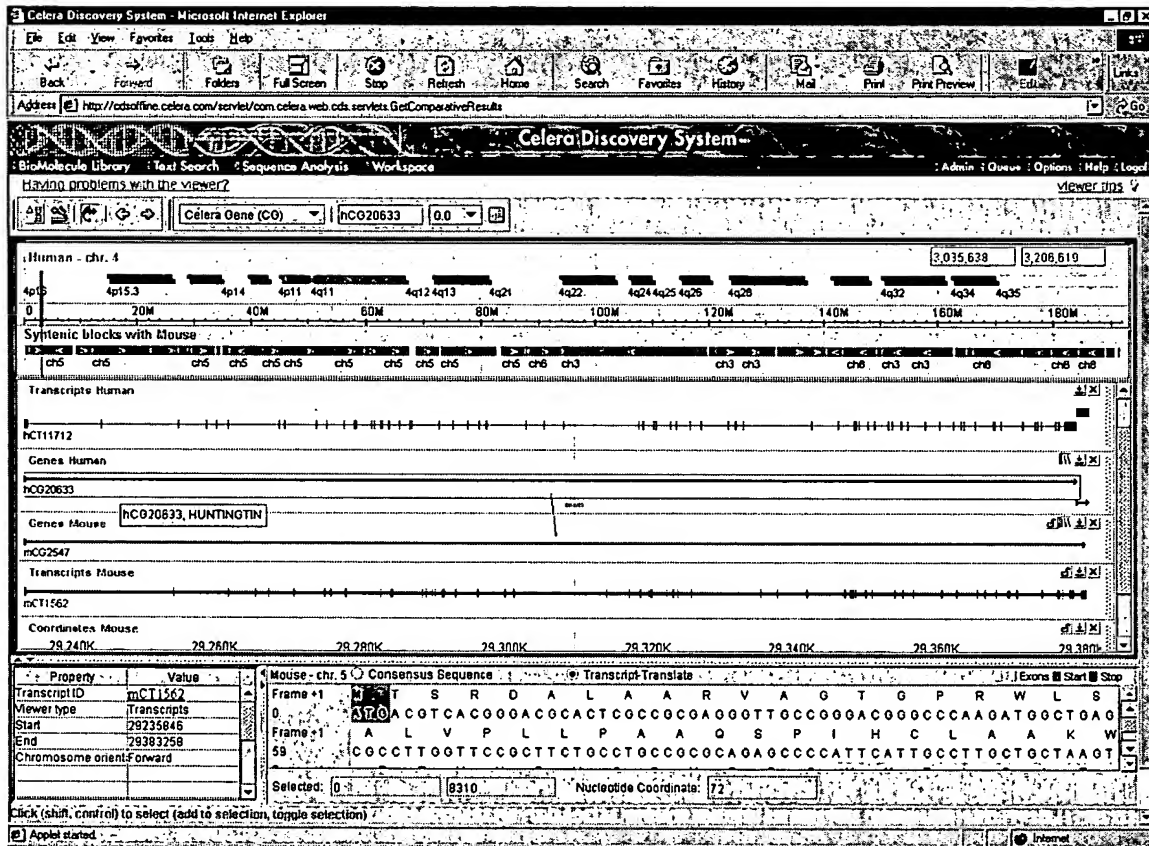


Figure - 24

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Inventors: Moore, et al

Atty. Ref.: 9692-000030

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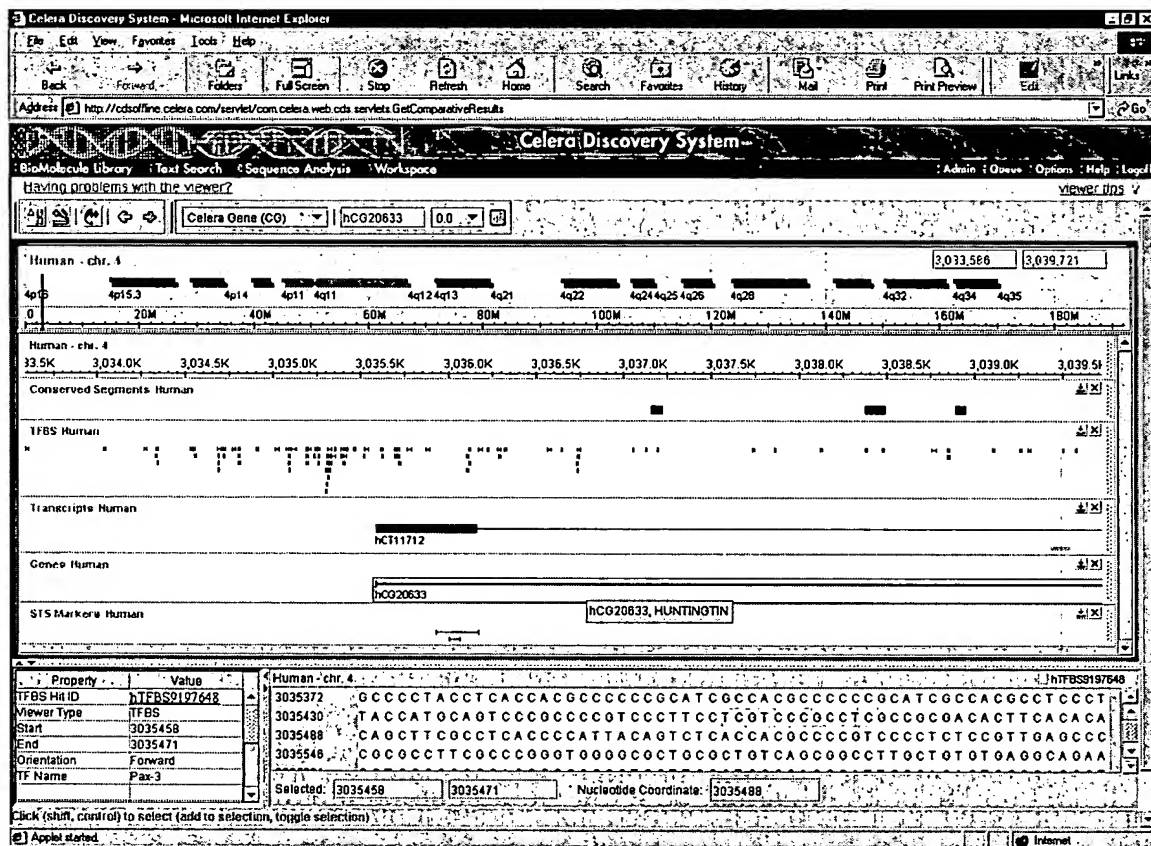


Figure - 25

Inventors: Moore, et al
Atty. Ref.: 9692-000030

Property	Value
Syntenic Block ID	hmSB1233
Viewer type	Syntenic Blocks
Start: Mouse	1149235
End: Mouse	3811691
Orientation: Mouse	Forward
Start: Human	1149235
End: Human	3811691
Chromosome: Mous	5
Chromosome: Hum	4
Orientation: Human	Forward

Property	Value
Transcript ID	hCT7498
Viewer type	Transcripts
Start	3254119
End	3400189
Chromosome orient	Forward

Property	Value
BAC ID	AL390065
Viewer type	Bac Clones
Start position	2876899
End position	3050577
Chromosome orient	Reverse

Property	Value
Conserved Segment	208000003331535
Viewer Type	Conserved Segme...
Start: Human	3991844
End: Human	3992014
Orientation: Human	Forward
Score	80
Chromosome: Mous	5
Begin: Mouse	32737628
End: Mouse	32737798
Orientation: Mouse	Forward

Property	Value
Cytoband Name	4p16
Viewer type	Cytobands
Start position	0
End position	15227048
Chromosome orient	Forward

Property	Value
STS Marker ID	UniSTS:7537
Viewer type	STS Markers
Start position	4029422
End position	4029563
Orientation	Uncertain

Property	Value
Scaffold ID	GA_xSJB7PYIMB
Viewer type	Scaffolds
Start position	4113701
End position	8846693
Chromosome orient	Reverse

Property	Value
Syntenic anchor ID	hmSA17902
Viewer type	Syntenic Anchors
Chromosome: Mous	5
Start: Mouse	2556830
End: Mouse	2557097
Orientation: Mouse	Forward
Chromosome: Hum	4
Start: Human	2556830
End: Human	2557097
Orientation: Human	Forward

Property	Value
TFBS Hit ID	hTFBS2211691
Viewer Type	TFBS
Start	1396467
End	1396485
Orientation	Forward
TF Name	NF-1

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